## **Amendments to the Specification**

Please replace the paragraph on page 6, lines 7-9 with the following amended paragraph:

FIG. 1 is a homology alignment of the amino acid sequences of the constant region of Cγ1 and Cγ3, aligned to maximize amino acid identity, and wherein non-conserved amino acids are identified by boxes. The amino acid sequences are presented in an N-terminal to C-terminal direction from left to right and top to bottom, spanning FIGS. 1A-1B. The Cγ1 amino acid sequence is labeled GC1/118 HUMAN, and is SEQ ID NO: 1. The Cγ3 amino acid sequence is labeled GC3 HUMAN/118, and is SEQ ID NO: 3;

Please replace the paragraph on page 6, lines 10-12 with the following amended paragraph:

FIG. 2 is a homology alignment of the amino acid sequences of constant region of Cγ1, Cγ2, and Cγ 4, aligned to maximize amino acid identity, and wherein non-conserved amino acids are identified by boxes. The amino acid sequences are presented in an N-terminal to C-terminal direction from left to right and top to bottom, spanning FIGS. 2A-2B. The Cγ1 amino acid sequence is labeled GC1/118 HUMAN, and is SEQ ID NO: 1. The Cγ2 amino acid sequence is labeled GC2/118 HUMAN, and is SEQ ID NO: 2. The Cγ4 amino acid sequence is labeled GC4/118 HUMAN, and is SEQ ID NO: 4;